## Fig. S1

>nbe-miR403a precursor

AAAGAGTCATATTTCACGTTTGTGCGTGATTCTGACAACCCTTTTTATCATTTTTATCATGTGGGTTGTGTTAGATTCACGCACAAACTCG CAATATGTCTTTTCT

Location: >Niben101Scf00793Ctg034:3501-4700

>nbe-miR403b precursor

**ACGCACAAACTCGTAATATGACTTTTCCTCATCT** 

Location: >Niben101Scf06412Ctg035:201-1400

>nbe-MIR162a precursor

TGGTTGGTTAAGAACACTGGAGGCAGCGGTT ATCGATCTGTTCCCTGAAAAGCGATAAACAAAATATAGCAACAGGAATCGG<mark>TCGATAAACC</mark>

**TCTGCATCCAG**CGGTTAACCCCCTTCT

Location: > Niben101Scf11303Ctg014:3301-4500

>nbe-MIR162b precursor

TTCTTTTGGTTGGTTAAAAACACTGGAGGCAGCGGTT ATCGATCTGTTCCCTGAAAGGCGATAACAAAAATATAGCAACAGGAATCGGTCGA

TAAACCTCTGCATCCAGCGCTTAACCCCCCTTCTCTCT

Location: >Niben101Scf01764Ctg011:2801-4000 (-1)

>nbe-MIR168a precursor

TGTCTCTAATTCGCTTGGTGCAGGTCGGGATTAATTCGCCGGCGACGGCGGCAATCACGACGACGGTGATTGTTATTTAATGGAGTTTAGAC

GTACGAAGTTATCAACATTTTTTTGTTTTGCAGCGAAATTTGTCCCGCCTTGCATCAACTGAATTGGAGACTGC

Location: >Niben101Scf10054Ctg018:1201-2400

>nbe-MIR168b precursor

. CTTGGTGCAGGTCGGGA<mark>T</mark>TAATTCGCCGGCGGCGGCAATCACGACGGCGGTGATTGTTATTAAATGGAGTTAAGATGTACGA

AGTTATCAACTTTTTTTGTTTTGCGGCGAAATTTGTCCCGCCTTGCATCAACTGAATTAGAGACTGC

Location: > Niben101Scf10307Ctg006:2101-3300

>nbe-MIR168c-1 precursor

GTTTATGACGAAGTTTGGGTCCCGCCTTGCATCAACTGAATCGGAGACTGCGGCGAAT

Location: >Niben101Scf04636Ctg025:701-1900

>nbe-MIR168c-2 precursor

TGTTGTTTATGACGAAGTTTGGGTCCCGCCTTGCATCAACTGAATCGGAGACTGCGGT

Location: >Niben101Scf07765Ctg021:44501-45800 (-1)

>nbe-MIR168d precursor

GGTCTCTAATTCGCTTGGTGCAGGTCGGGA\_CTGACTCGCCGGTGACGCTGTCCGCGCCGGAATCGGCGCCCTTAGTTTCATTGATATTTTG TTTACATGTACCGATGTTTAGAGAGCTCTCTTTTGGACTTATTTGTTAGTAGTTTTTTGTTGTTGAAAATAACTTGATATATGTATAGCTAAGTAGA 

TTAATAGATTAGTTATGAATGAAAGTGCGCCTCGTAAATGGCTAAGTTTGTTCCCGCCTTGCATCAACTGAATTGGTGACCGC

Location: >Niben101Scf08513Ctg014:21401-22800 (-1)

>nbe-MIR168e precursor

TTCAATTTTTTTAATAGATTAGTTATGAATGAAATTGCGCCTCGTCAATGGCTAAGTTTGTTCCCGCCTTGCATCAACTGAATCGGTGACCGC

Location: >Niben101Scf07340Ctg008:19901-21400

Fig. S1. Nicotiana benthamiana miRNA precursor 5'-to-3' sequences; their location in

the Nb genome v.0.4.2 at <a href="https://solgenomics.net/">https://solgenomics.net/</a>. Mature and passenger sequences

are highlighted in red (mature) and in blue (passenger strands); nucleotides that differ

between miRNA isoforms are labeled in green. The search was supported by unpublished

sRNA data and facilitated by the fact that miRNA guide strands are highly conserved among

dicots (44).